

MLLNVLRICI	IVCLVNDGAG	KHSEGRERTK	TYSLNSRGYF	40
RKERGARRSK	ILLVNTKGLD	EPHIGHGDFG	LVAELFDSTR	80
THTNRKEPDM	NKVKLFSTVA	HG <u>NKS</u> ARRKA	<u>YNGS</u> RRNIFS	120
RRSFDKRNTE	VTEKPGAKMF	WNNFLVKMNG	APONT <u>SH</u> GSK	160
AQEIMKEACK	TLPFTQNIVH	ENCDRMVION	NLCFGKCISL	200
HVPNQQDRRN	TCSHCLPSKF	TLNHL <u>TLNCT</u>	GSKNVVKVVM	240
MVEECTCEAH	KSNFHQTAQF	NMDTSTTLHH		270

Figure 1. Deduced amino acid sequence of *Xenopus cerberus* protein. SEQ ID NO:1.

Figure 2. Nucleotide sequence of the full-length cerberus DNA derived from the *Xenopus* organizer. The sense strand is on top (in the 5' to 3' direction) and the antisense strand on the bottom line (on the opposite direction). SEQ ID NO:2.

GAATTCCCAG	CAAGTCGCTC	AGAAACACTG	CAGGGTCTAG	ATATCATACA	ATGTTACTAA	60
CTTAAGGGTC	GTTCAGCGAG	TCTTTGTGAC	GTCCAGATC	TATAGTATGT	TACAATGATT	
ATGTACTCAG	GATCTGTATT	ATCGTCTGCC	TTGTGAATGA	TGGAGCAGGA	AAACACTCAG	120
TACATGAGTC	CTAGACATAA	TAGCAGACGG	AACACTTACT	ACCTCGTCCT	TTTGTGAGTC	
AAGGACGAGA	AAGGACAAAA	ACATATTAC	TTAACAGCAG	AGGTACTTTC	AGAAAAGAAA	180
TTCTGCTCT	TTCTGTTTT	TGTATAAGTG	AATTGTCGTC	TCCAATGAAG	TCTTTTCTTT	
GAGGAGCAG	TAGGAGCAAG	ATTCTGCTGG	TGAATACTAA	AGGTCTTGAT	GAACCCCA	240
CTCTCGTGC	ATCCTCGTTC	TAAGACGACC	ACTTATGATT	TCCAGAATA	CTGGGGTGT	
TTGGGCATGG	TGATTTTCGC	TTAGTAGCTG	AACATTTGA	TTCCACCAGA	ACACATACAA	300
AACCCGTACC	ACTAAAAGCG	AATCATCGAC	TTGATAAACT	AAGGTGGTCT	TGTGTATGTT	
ACAGAAAAGA	GCCAGACATG	AACAAAGTCA	AGCTTTTCTC	AACAGTTGCC	CATGGAAACA	360
TGTCTTTTCT	CGGTCTGTAC	TTGTTTCAGT	TCGAAAAGAG	TTGTCAACGG	GTACCTTTGT	
AAAGTGCAAG	AAGAAAAGCT	TACAATGGTT	CTAGAAGGAA	TATTTTTCCT	CGCCGTTCTT	420
TTTCACGTTT	TTCTTTTCGA	ATGTTACCAA	GATCTTCCTT	ATAAAAAGGA	GCGGCAAGAA	
TTGATAAAG	AAATACAGAG	GTTACTGAAA	AGCCTGGTGC	CAAGATGTTT	TGGAACAATT	480
AACTATTTTC	TTTATGTCTC	CAATGACTTT	TCGGACCACG	GTTCTACAAG	ACCTTGTTAA	
TTTTGGTTAA	AATGAATGGA	GCCCCACAGA	ATACAAGCCA	TGGCAGTAAA	GCACAGGAAA	540
AAAACCAATT	TTACTTACCT	CGGGGTGTCT	TATGTTCCGG	ACCGTCATTT	CGTGTCTTTT	
TAATGAAAGA	AGCTTGCAAA	ACCTTGTTTT	TCCTCAGAA	TATTGTACAT	GAAACTGTG	600
ATTACTTTCT	TCGAACGTTT	TGGAACAAAA	AGTGAGTCTT	ATAACATGTA	CTTTTGACAC	
ACAGGATGGT	GATACAGAAC	AATCTGTGCT	TTGGTAAATG	CATCTCTCTC	CATGTTCCAA	660
TGTCCTACCA	CTATGTCTTG	TTAGACACGA	AACCATTAC	GTAGAGAGAG	GTACAAGGTT	
ATCAGCAAGA	TCGACGAAAT	ACTTGTTCCC	ATTGCTTGCC	GTCCAAATTT	ACCCTGAACC	720
TAGTCGTTCT	AGCTGCTTTA	TGAACAAGGG	TAACGAACGG	CAGGTTTAAA	TGGGACTTGG	
ACCTGACGCT	GAATTGTACT	GGATCTAAGA	ATGTAGTAAA	GGTTGTCATG	ATGGTAGAGG	780
TGGACTGCGA	CTTAACATGA	CCTAGATTCT	TACATCATTT	CCAACAGTAC	TACCATCTCC	
AATGCACGTG	TGAAGCTCAT	AAGAGCAACT	TCCACCAAAC	TGCACAGTTT	AACATGGATA	840
TTACGTGCAC	ACTTCGAGTA	TTCTCGTTGA	AGGTGGTTTG	ACGTGTCAAA	TTGTACCTAT	
CATCTACTAC	CCTGCACCAT	TAAAGGACTG	CCATACAGTA	TGGAAATGCC	CTTTTGTGG	900
GTAGATGATG	GGACGTGGTA	ATTCCTGAC	GGTATGTCAT	ACCTTTACGG	GAAAACAACC	
AATATTTGTT	ACATACTATG	CATCTAAAGC	ATTATGTTGC	CTTCTATTTT	ATATAACCAC	960
TTATAAACAA	TGTATGATAC	GTAGATTTTG	TAATACAACG	GAAGATAAAG	TATATTGGTG	
ATGGAATAAG	GATTGTATGA	ATTATAATTA	ACAAATGGCA	TTTTGTGTAA	CATGCAAGAT	1020
TACCTTATTC	CTAACATACT	TAATATTAAT	TGTTTACCGT	AAAACACATT	GTACGTTCTA	

CTCTGTTCCA	TCAGTTGCAA	GATAAAAGGC	AATATTTGTT	TGACTTTTTT	TCTACAAAAT	1080
GAGACAAGGT	AGTCAACGTT	CTATTTTCCG	TTATAAACAA	ACTGAAAAAA	AGATGTTTTA	
GAATACCCAA	ATATATGATA	AGATAATGGG	GTCAAAACTG	TTAAGGGGTA	ATGTAATAAT	1140
CTTATGGGTT	TATATACTAT	TCTATTACCC	CAGTTTTGAC	AATTCCCAT	TACATTATTA	
AGGGACTAAG	TTTGCCCAGG	AGCAGTGACC	CATAACAACC	AATCAGCAGG	TATGATTTAC	1200
TCCCTGATTG	AAACGGGTCC	TCGTCACTGG	GTATTGTTGG	TTAGTCGTCC	ATACTAAATG	
TGGTCACCTG	TTTAAAAGCA	AACATCTTAT	TGGTTGCTAT	GGGTTACTGC	TTCTGGGCAA	1260
ACCAGTGGAC	AAATTTTCGT	TTGTAGAATA	ACCAACGATA	CCCAATGACG	AAGACCCGTT	
AATGTGTGCC	TCATAGGGGG	GTTAGTGTGT	TGTGTACTGA	ATAAATTGTA	TTTATTTTCAT	1320
TTACACACGG	AGTATCCCCC	CAATCACACA	ACACATGACT	TATTTAACAT	AAATAAAGTA	
TGTTACAAAA	AAAAAAAA					
ACAATGTTTT	TTTTTTTT					

Fig. 2. (Continuation page 2, SEQ ID NO:2).

MSRTRKVDL LLLAIPGLAL LLLPNAYCAS CEPVRIPMCK SMPWNMTKMP NHLHHSTQAN 60
AILAIEQFEG LLTTECSQDL LFFLCAMYAP ICTIDFQHEP IKPCKSV CER ARAGCEPILI 120
KYRHTWPESL ACEELPVYDR GVCISPEAIV TVEQGTDSMP DFSMDSNNGN CGSGREHCKC 180
KPMKATQKTY LKNYNYVIR AKVKEVKVVC HDATAIVEVK EILKSSLVNI PKDTVTLYTN 240
SGCLCPQLVA NEEYIIMGYE DKERTRLLLV EGSLAEKWRD RLAKKVKRWD QKLRRPRKSK 300
DPVAPIPNKN SNSRQARS

Figure 3. Deduced amino acid sequence of Xenopus frazzled protein. SEQ ID NO:3.

Figure 4. Nucleotide sequence of the full-length frazzled cDNA derived from the *Xenopus* organizer. The sense strand of the DNA on top (5' to 3' direction) and the antisense strand on the bottom line (opposite direction). SEQ ID NO:4.

GAATTCCTT TCACACAGGA CTCCTGGCAG AGGTGAATGG TTAGCCCTAT GGATTGTT	60
CTTAAGGGAA AGTGTGTCCT GAGGACCGTC TCCACTTACC AATCGGGATA CCTAAACCAA	
TGTTGATTTT GACACATGAT TGATTGCTTT CAGATAGGAT TGAAGGACTT GGATTTTAT	120
ACAATAAAA CTGTGTACTA ACTAACGAAA GTCTATCCTA ACTTCCTGAA CCTAAAAATA	
CTAATCTGCG ACTTTTAAAT TATCTGAGTA ATTGTTCAAT TTGTATTGGA TGGGACTAAA	180
GATTAAGACG TGAAAAATTA ATAGACTCAT TAACAAGTAA AACATAACCT ACCCTGATTT	
GATAAACTTA ACTCCTTGCT TTTGACTTGC CCATAAACTA TAAGGTGGGG TGAGTTGTAG	240
CTATTTGAAT TGAGGAACGA AAACGAAACG GGTATTTGAT ATTCCACCCC ACTCAACATC	
TTGCTTTTAC ATGTGCCCGAG ATTTTCCCTG TATTCCTGT ATTCCCTCTA AAGTAAGCCT	300
AACGAAAATG TACACGGGTC TAAAGGGAC ATAAGGGACA TAAGGGAGAT TTCATTCCGA	
ACACATACAG GTTGGGCAGA ATAACAATGT CTCGAACAAG GAAAGTGGAC TCATTACTGC	360
TGTGTATGTC CAACCCGTCT TATTGTTACA GAGCTTGTTT CTTTCACCTG AGTAATGACG	
TACTGGCCAT ACCTGGACTG GCGCTTCTCT TATTACCCAA TGCTTACTGT GCTTCGTGTG	420
ATGACCGGTA TGGACCTGAC CGCGAAGAGA ATAATGGGTT ACGAATGACA CGAAGCACAC	
AGCCTGTGCG GATCCCCATG TGCAAATCTA TGCCATGGAA CATGACCAAG ATGCCCAACC	480
TCGGACACGC CTAGGGGTAC ACGTTTAGAT ACGGTACCTT GTACTGGTTC TACGGGTTGG	
ATCTCCACCA CAGCACTCAA GCCAATGCCA TCCTGGCAAT TGAACAGTTT GAAGGTTTGC	540
TAGAGGTGGT GTCGTGAGTT CGGTTACGGT AGGACCGTTA ACTTGTCAA CTTCCAAACG	
TGACCACTGA ATGTAGCCAG GACCTTTTGT TCTTCTGTG TGCCATGTAT GCCCCCATTT	600
ACTGGTGAAT TACATCGGTC CTGGAAAACA AGAAAGACAC ACGGTACATA CGGGGGTAAA	
GTACCATCGA TTTCCAGCAT GAACCAATTA AGCCTTGCAA GTCCGTGTGC GAAAGGGCCA	660
CATGGTAGCT AAAGGTCGTA CTTGGTTAAT TCGGAACGTT CAGGCACACG CTTTCCCGGT	
GGGCCGGCTG TGAGCCCATT CTCATAAAGT ACCGGCACAC TTGGCCAGAG AGCCTGGCAT	720
CCCGGCCGAC ACTCGGGTAA GAGTATTTCA TGGCCGTGTG AACCGGTCTC TCGGACCGTA	
GTGAAGAGCT GCGCGTATAT GACAGAGGAG TCTGCATCTC CCCAGAGGCT ATCGTCACAG	780
CACTTCTCGA CGGGCATATA CTGTCTCCTC AGACGTAGAG GGGTCTCCGA TAGCAGTGTC	
TGGAACAAGG AACAGATTCA ATGCCAGACT TCTCCATGGA TTCAAACAAT GGAAATTGCG	840
ACCTTGTTCC TTGTCTAAGT TACGGTCTGA AGAGGTACCT AAGTTTGTTA CCTTTAACGC	
GAAGCGGCAG GGAGCACTGT AAATGCAAGC CCATGAAGGC AACCCAAAAG ACGTATCTCA	900
CTTCGCCGTC CCTCGTGACA TTTACGTTCC GGTACTTCCG TTGGGTTTTT TGCATAGAGT	
AGAATAATTA CAATTATGTA ATCAGAGCAA AAGTGAAAGA GGTGAAAGTG AAATGCCACG	960
TCTTATTAAT GTTAATACAT TAGTCTCGTT TTCACTTTCT CCACTTTCAC TTTACGGTGC	
ACGCAACAGC AATTGTGGAA GTAAAGGAGA TTCTCAAGTC TTCCCTAGTG AACATTCTTA	1020
TGCGTTGTGCG TTAACACCTT CATTTCTCTT AAGAGTTCAG AAGGGATCAC TTGTAAGGAT	

AAGACACAGT	GACACTGTAC	ACCAACTCAG	GCTGCTTGTG	CCCCCAGCTT	GTTGCCAATG	1080
TTCTGTGTCA	CTGTGACATG	TGGTTGAGTC	CGACGAACAC	GGGGGTGAA	CAACGGTTAC	
AGGAATACAT	AATTATGGGC	TATGAAGACA	AAGAGCGTAC	CAGGCTTCTA	CTAGTGGAAG	1140
TCCTTATGTA	TTAATACCCG	ATACTTCTGT	TTCTCGCATG	GTCCGAAGAT	GATCACCTTC	
GATCCTTGGC	CGAAAAATGG	AGAGATCGTC	TTGCTAAGAA	AGTCAAGCGC	TGGGATCAAA	1200
CTAGGAACCG	GCTTTTTACC	TCTCTAGCAG	AACGATTCTT	TCAGTTCGCG	ACCCTAGTTT	
AGCTTCGACG	TCCCAGGAAA	AGCAAAGACC	CCGTGGCTCC	AATTCCCAAC	AAAAACAGCA	1260
TCGAAGCTGC	AGGGTCCTTT	TCGTTTCTGG	GGCACCAGG	TTAAGGGTTG	TTTTTGTCGT	
ATTCCAGACA	AGCGCGTAGT	TAGACTAACG	GAAAGGTGTA	TGGAACTCT	ATGGACTTTG	1320
TAAGGTCTGT	TCGCGCATCA	ATCTGATTGC	CTTTCCACAT	ACCTTTGAGA	TACCTGAAAC	
AAACTAAGAT	TTGCATTGTT	GGAAGAGCAA	AAAAGAAATT	GCACTACAGC	ACGTTATATT	1380
TTTGATTCTA	AACGTAACAA	CCTTCTCGTT	TTTTCTTTAA	CGTGATGTCG	TGCAATATAA	
CTATTGTTTA	CTACAAGAAG	CTGGTTTAGT	TGATTGTAGT	TCTCCTTTCC	TTCTTTTTTT	1440
GATAACAAAT	GATGTTCTTC	GACCAAATCA	ACTAACATCA	AGAGGAAAGG	AAGAAAAAAA	
TTATACTAT	ATTTGCACGT	GTTCCCAGGC	AATTGTTTTA	TTCAACTTCC	AGTGACAGAG	1500
AATATTGATA	TAAACGTGCA	CAAGGGTCCG	TTAACAAAT	AAGTTGAAGG	TCACTGTCTC	
CAGTGACTGA	ATGTCTCAGC	CTAAGAAGC	TCAATTCATT	TCTGATCAAC	TAATGGTGAC	1560
GTCAGTACT	TACAGAGTCG	GATTTCTTCG	AGTTAAGTAA	AGACTAGTTG	ATTACCACTG	
AAGTGTTTGA	TACTTGGGGA	AAGTGAACCT	ATTGCAATGG	TAAATCAGAG	AAAAGTTGAC	1620
TTCACAACT	ATGAACCCCT	TTCATTGAT	TAACGTTACC	ATTTAGTCTC	TTTCAACTG	
CAATGTTGCT	TTTCCTGTAG	ATGAACAAGT	GAGAGATCAC	ATTTAAATGA	TGATCACTTT	1680
GTTACAACGA	AAAGGACATC	TACTTGTTCA	CTCTCTAGTG	TAAATTTACT	ACTAGTGAAA	
CCATTTAATA	CTTTCAGCAG	TTTTAGTTAG	ATGACATGTA	GGATGCACCT	AAATCTAAAT	1740
GGTAAATTAT	GAAAGTCGTC	AAAATCAATC	TACTGTACAT	CCTACGTGGA	TTTAGATTTA	
ATTTTATCAT	AAATGAAGAG	CTGGTTTAGA	CTGTATGGTC	ACTGTTGGGA	AGGTAAATGC	1800
TAAAATAGTA	TTTACTTCTC	GACCAATCT	GACATACCAG	TGACAACCCCT	TCCATTTACG	
CTACTTTGTC	AATTCTGTTT	TAAAAATTGC	CTAAATAAAT	ATTAAGTCCT	AAATAAAAAA	1860
GATGAAACAG	TTAAGACAAA	ATTTTAAACG	GATTTATTTA	TAATTCAGGA	TTTATTTTTT	
AAAAAAAAAA	AAAAA					
TTTTTTTTTT	TTTTT					

Fig. 4. (Continuation page 2, SEQ ID NO:4).

MLLLFRAIPM LLLGLMVLQT DCEIAQYYID EEEPPGTVIA VLSQHSIFNT TDIPATNFRL	60
MKQFNNSLIG VRESQGQLSI MERIDREQIC RQSLHCNLAL DVVSFSKGHF KLLNVKVEVR	120
DINDHSPHFP SEIMHVEVSE SSSVGTRIPL EIAIDEDVGS NSIQNFQISN NSHFSIDVLT	180
RADGVKYADL VLMRELDREI QPTYIMELLA MDGGVPSLSG TAVVNIRVLD FNDNSPVFER	240
STIAVDLVED APLGYLLEL HATDDDEGVN GEIVYGFSTL ASQEVRLFK INSRTGSVTL	300
EGQVDFETKQ TYEFEVQAQD LGPNPLTATC KVTVHILDVN DNTPAITITP LTTVNAGVAY	360
IPETATKENF IALISTTDRA SGSNGQVRCT LYGHEHFKLQ QAYEDSYMIV TTSTLDRENI	420
AAYSLTVVAE DLGFPSLGTK KYITVKVSD E NDNA PVFSKP QYEASILENN APGSYITTVI	480
ARDSDSQNG KVNRYRLVDAK VMQSLTTFV SLDADSGVLR AVRSLDYEKL KQLDFEIEAA	540
DNGIPQLSTR VQLNLRIVDQ NDNCPVITNP LLNNGSGEVL LPISAPQNYL VFQLKAEDSD	600
EGHNSQLFYT ILRDPSRLFA INKESGEVFL KKQLNSDHSE DLSIVVAVYD LGRPSLSTNA	660
TVKFILTDSF PSNVEVVILQ PSAEEQHQID MSIIFIAVLA GGCALLLLAI FVFACTCKKK	720
AGEFKQVPEQ HGTCNEERLL STPSPQSVSS SLSQSESCQL SINTESENC VSSNQEQHQQ	780
TGIKHSISVP SYHTSGWHL D NCAM SISGHS HMGHISTKVQ WAKEIVTSMT VTLILVENQK	840
RRALSSQCRH KPVLTNTQMNQ QGSDMPITIS ATESTRVQKM GTAHCNMKRA IDCLTL	

Figure 5. Deduced amino acid sequence of the *Xenopus* PAPC (paraxial protocadherin) protein. It encodes a member of the cadherin family of transmembrane proteins that has dorsalizing activity when constructs are injected into *Xenopus* embryos. SEQ ID NO:5.

Figure 6. Nucleotide sequence of the full-length PAPC cDNA derived from the *Xenopus* organizer. The sense strand of the DNA is shown in the top line (in the 5' to 3' direction), and the bottom line shows the antisense strand (opposite orientation). SEQ ID NO:6.

GAATTCCCAG	AGATGAACTC	CTTGAGATTG	TTTTAAATGA	CTGCAGGTCT	GGAAGGATTC	60
CTTAAGGGTC	TCTACTTGAG	GAACTCTAAC	AAAATTTACT	GACGTCCAGA	CCTTCCTAAG	
ACATTGCCAC	ACTGTTTCTA	GGCATGAAAA	AACTGCAAGT	TTCAACTTTG	TTTTTGGTGC	120
TGTAACGGTG	TGACAAAGAT	CCGTACTTTT	TTGACGTTCA	AAGTTGAAAC	AAAAACCACG	
AACTTTGATT	CTTCAAGATG	CTGCTTCTCT	TCAGAGCCAT	TCCAATGCTG	CTGTTGGGAC	180
TTGAAACTAA	GAAGTTCTAC	GACGAAGAGA	AGTCTCGGTA	AGGTTACGAC	GACAACCCCTG	
TGATGGTTTT	ACAAACAGAC	TGTGAAATTG	CCCAGTACTA	CATAGATGAA	GAAGAACCCC	240
ACTACCAAAA	TGTTTGTCTG	ACACTTTAAC	GGGTCATGAT	GTATCTACTT	CTTCTTGGGG	
CTGGCACTGT	AATTGCAGTG	TTGTCACAAC	ACTCCATATT	TAACACTACA	GATATACCTG	300
GACCGTGACA	TTAACGTCAC	AACAGTGTG	TGAGGTATAA	ATTGTGATGT	CTATATGGAC	
CAACCAATTT	CCGTCTAATG	AAGCAATTTA	ATAATTCCTT	TATCGGAGTC	CGTGAGAGTG	360
GTTGGTTAAA	GGCAGATTAC	TTCGTTAAAT	TATTAAGGGA	ATAGCCTCAG	GCACTCTCAC	
ATGGGCAGCT	GAGCATCATG	GAGAGGATTG	ACCGGGAGCA	AATCTGCAGG	CAGTCCCTTC	420
TACCCGTCGA	CTCGTAGTAC	CTCTCCTAAC	TGGCCCTCGT	TTAGACGTCC	GTCAGGGAAG	
ACTGCAACCT	GGCTTTGGAT	GTGGTCAGCT	TTTCCAAAGG	ACACTTCAAG	CTTCTGAACG	480
TGACGTTGGA	CCGAAACCTA	CACCACTCGA	AAAGGTTTCC	TGTGAAGTTC	GAAGACTTGC	
TGAAAGTGGA	GGTGAGAGAC	ATTAATGACC	ATAGCCCTCA	CTTTCCCAGT	GAAATAATGC	540
ACTTTCACCT	CCACTCTCTG	TAATTACTGG	TATCGGGAGT	GAAAGGGTCA	CTTTATTACG	
ATGTGGAGGT	GTCTGAAAGT	TCCTCTGTGG	GCACCAGGAT	TCCTTTAGAA	ATTGCAATAG	600
TACACCTCCA	CAGACTTTCA	AGGAGACACC	CGTGGTCCTA	AGGAAATCTT	TAACGTTATC	
ATGAAGATGT	TGGGTCCAAC	TCCATCCAGA	ACTTTCAGAT	CTCAAATAAT	AGCCACTTCA	660
TACTTCTACA	ACCCAGGTTG	AGGTAGGTCT	TGAAAGTCTA	GAGTTTATTA	TCGGTGAAGT	
GCATTGATGT	GCTAACCAGA	GCAGATGGGG	TGAAATATGC	AGATTTAGTC	TTAATGAGAG	720
CGTAACTACA	CGATTGGTCT	CGTCTACCCC	ACTTTATACG	TCTAAATCAG	AATTACTCTC	
AACTGGACAG	GGAAATCCAG	CCAACATACA	TAATGGAGCT	ACTAGCAATG	GATGGGGGTG	780
TTGACCTGTC	CCTTTAGGTC	GGTTGTATGT	ATTACCTCGA	TGATCGTTAC	CTACCCCCAC	
TACCATCACT	ATCTGGTACT	GCAGTGTTTA	ACATCCGAGT	CCTGGACTTT	AATGATAACA	840
ATGGTAGTGA	TAGACCATGA	CGTCACCAAT	TGTAGGCTCA	GGACCTGAAA	TTACTATTGT	
GCCCAGTGTT	TGAGAGAAGC	ACCATTGCTG	TGGACCTAGT	AGAGGATGCT	CCTCTGGGAT	900
CGGGTCACAA	ACTCTCTTCG	TGGTAACGAC	ACCTGGATCA	TCTCCTACGA	GGAGACCCTA	
ACCTTTTGTT	GGAGTTACAT	GCTACTGACG	ATGATGAAGG	AGTGAATGGA	GAAATTGTTT	960
TGGAAACAA	CCTCAATGTA	CGATGACTGC	TACTACTTCC	TCACTTACCT	CTTTAACAAA	
ATGGATTGAG	CACCTTTGGC	TCTCAAGAGG	TACGTGAGCT	ATTTAAAATT	AACTCCAGAA	1020
TACCTAAGTC	GTGAAACCGT	AGAGTTCTCC	ATGCAGTCGA	TAAATTTTAA	TTGAGGTCTT	

AAGTCGTTAT TTTGCAACCA TCTGCAGAAG AGCAGCACCA GATCGATATG TCCATTATAT	2220
TTTCAGCAATA AAACGTTGGT AGACGTCTTC TCGTCGTGGT CTAGCTATAC AGGTAATATA	
TCATTGCAGT GCTGGCTGGT GGTGTGCTT TGCTACTTTT GGCCATCTTT TTTGTGGCCT	2280
AGTAACGTCA CGACCGACCA CCAACACGAA ACGATGAAAA CCGGTAGAAA AAACACCGGA	
GTAATTGTAA AAAGAAAGCT GGTGAATTTA AGCAGGTACC TGAACAACAC GGAACATGCA	2340
CATGAACATT TTTCTTTCGA CCACTTAAAT TCGTCCATGG ACTTGTGTG CCTTGTACGT	
ATGAAGAACG CCTGTTAAGC ACCCCATCTC CCCAGTCGGT CTCTTCTTCT TTGTCTCAGT	2400
TACTTCTTGC GGACAATTGC TGGGGTAGAG GGGTCAGCCA GAGAAGAAGA AACAGAGTCA	
CTGAGTCATG CCAACTCTCC ATCAATACTG AATCTGAGAA TTGCAGCGTG TCCTCTAACC	2460
GACTCAGTAC GGTGAGAGG TAGTTATGAC TTAGACTCTT AACGTCGCAC AGGAGATTGG	
AAGAGCAGCA TCAGCAAACA GGCATAAAGC ACTCCATCTC TGTACCATCT TATCACACAT	2520
TTCTCGTCGT AGTCGTTTGT CCGTATTTTCG TGAGGTAGAG ACATGGTAGA ATAGTGTGTA	
CTGGTTGGCA CCTGGACAAT TGTGCAATGA GCATAAGTGG ACATTCTCAC ATGGGGCACA	2580
GACCAACCGT GGACCTGTTA ACACGTTACT CGTATTCACC TGTAAGAGTG TACCCCGTGT	
TTAGTACAAA GGTACAGTGG GCAAAGGAGA TAGTGACTTC AATGACAGTG ACTCTGATAC	2640
AATCATGTTT CCATGTCACC CGTTTCTCT ATCACTGAAG TTAGTGTAC TGAGACTATG	
TAGTGGAGAA TCAGAAAAGA AGAGCATTGA GCAGCCAATG CAGGCACAAG CCAGTGCTCA	2700
ATCACCTCTT AGTCTTTTCT TCTCGTAACT CGTCGGTTAC GTCCGTGTTT GGTACAGAGT	
ATACACAGAT GAATCAGCAG GGTCCGACA TGCCGATAAC TATTTAGCC ACCGAATCAA	2760
TATGTGTCTA CTTAGTCGTC CCAAGGCTGT ACGGCTATTG ATAAAGTCGG TGGCTTAGTT	
CAAGGGTCCA GAAAATGGGA ACTGCACATT GCAATATGAA AAGGGCTATA GACTGTCTTA	2820
GTTCCAGGT CTTTTACCCT TGACGTGTAA CGTTATACTT TTCCCGATAT CTGACAGAAT	
CTCTGTAGCT CCTGTATATT ACAATACCTA CCATGCAAGA ATGCCTAACC TGCACATACC	2880
GAGACATCGA GGACATATAA TGTTATGGAT GGTACGTTCT TACGGATTGG ACGTGTATGG	
GAACCATACC CTTAGAGACC CTTATTACCA TATCAATAAT CCTGTTGCTA ATCGGATGCA	2940
CTTGGTATGG GAATCTCTGG GAATAATGGT ATAGTTATTA GGACAACGAT TAGCCTACGT	
GGCGGAATAT GAAAGAGATT TAGTCAACAG AAGTGCAACG TTATCTCCGC AGAGATCGTC	3000
CCGCCTTATA CTTTCTCTAA ATCAGTTGTC TTCACGTTGC AATAGAGGCG TCTCTAGCAG	
TAGCAGATAC CAAGAATTCA ATTACAGTCC GCAGATATCA AGACAGCTTC ATCCTTCAGA	3060
ATCGTCTATG GTTCTTAAGT TAATGTCAGG CGTCTATAGT TCTGTGGAAG TAGGAAGTCT	
AATTGCTACA ACCTTTTAAT CATTAGGCAT GCAAGTGAGA ATGCACAAAG GCAAGTGCTT	3120
TTAACGATGT TGGAAAATTA GTAATCCGTA CGTTCACTCT TACGTGTTTC CGTTCACGAA	
TAGCATGAAA GCTAAATATA TGGAGTCTCC CCTTTCCCTC TGATGGATGG GGGGAGACAC	3180
ATCGTACTTT CGATTTATAT ACCTCAGAGG GGAAAGGGAG ACTACCTACC CCCCTCTGTG	
AGGACAGTGC ATAAATATAC AGCTGCTTTC TATTTGCATT TCACTTGGGA ATTTTTGTG	3240
TCCTGTCACG TATTTATATG TCGACGAAAG ATAAACGTAA AGTGAACCTT TAAAAACAA	
TTTTTTACAT ATTTATTTTT CCTGAATTGA ATGTGACATT GTCCTGTCAC CTAAGTAGCA	3300
AAAAAATGTA TAAATAAAAA GGACTTAACT TACACTGTAA CAGGACAGTG GATTGATCGT	

Fig. 6. (Continuation page 3, SEQ ID NO:6).

ATTAAATCCA CAGACCTACA GTCAAATATT TGAGGGCCCC TGAAACAGCA CATCAGTCAG	3360
TAATTTAGGT GTCTGGATGT CAGTTTATAA ACTCCCGGGG ACTTTGTCGT GTAGTCAGTC	
GACCTAAAGT GGCCTTTTTA CTTTTCAGCAG CTCCTGGGTC TGCCCTCTGT GTTAATCAGC	3420
CTGGATTTC ACGGAAAAAT GAAAATCGTC GAGGACCCAG ACGGGAGACA CAATTAGTCG	
CCCTGGTCAA GTCCTGAGTA GGATCATGGC GTTTTATAT GCATCTCACC TACTTTGGAC	3480
GGGACCAGTT CAGGACTCAT CCTAGTACCG CAAAAATATA CGTAGAGTGG ATGAAACCTG	
GTGATTTACA CATAATAGGA AACGCTTGGT TTCAGTGAAG TCTGTGTTGT ATATATTCTG	3540
CACTAAATGT GTATTATCCT TTGCGAACCA AAGTCACTTC AGACACAACA TATATAAGAC	
TTATATACAC GCATTTTGTG TTTGTGTATA TATTTCAAGT CCATTGAGAT ATGTGTATAT	3600
AATATATGTG CGTAAACAC AAACACATAT ATAAAGTTCA GGTAAGTCTA TACACATATA	
AGTGCAGACC TTGTAAATTA AATATTCTGA TACTTTTCC TCAATAAATA TTAAAT	
TCACGTCTGG AACATTTAAT TTATAAGACT ATGAAAAGG AGTTATTAT AAATTTA	

Fig. 6. (Continuation page 4, SEQ ID NO:6).

MVCCGPGRML LGWAGLLVLA ALCLLQVPGA QAAACEPVRI PLCKSLPWNM TKMPNHLHHS 60
TQANAILAME QFEGLLGTHC SPDLLFFLCA MYAPICTIDF QHEPIKPCKS VCERARQGCE 120
PILIKYRHSW PESLACDELP VYDRGVCISP EAIVTADGAD FPMDSSTGHC RGASSERCKC 180
KPVRATQKTY FRNNYNYVIR AKVKEVKMKC HDVTAVVEVK EILKASLVNI PRDTVNLYTT 240
SGCLCPPLTV NEEYVIMGYE DEERSRLLLV EGSIAEKWKD RLGKKVKRWD MKLRHLGLGK 300
TDASDSTQNQ KSGRNSNPRP ARS.

Figure 7. Deduced amino acid sequence of mouse FRZB-1 protein. SEQ ID NO:7.

Figure 8. Nucleotide sequence of the full-length mouse FRZB-1 cDNA. SEQ ID NO:8.

AAGCCTGGGA	CCATGGTCTG	CTGCGGCCCG	GGACGGATGC	TGCTAGGATG	GGCCGGGTTG	60
TTCGGACCCCT	GGTACCAGAC	GACGCCGGGC	CCTGCCTACG	ACGATCCTAC	CCGGCCCAAC	
CTAGTCCTGG	CTGCTCTCTG	CCTGCTCCAG	GTGCCCCGAG	CTCAGGCTGC	AGCCTGTGAG	120
GATCAGGACC	GACGAGAGAC	GGACGAGGTC	CACGGGCCTC	GAGTCCGACG	TCGGACACTC	
CCTGTCCGCA	TCCCGCTGTG	CAAGTCCCTT	CCCTGGAACA	TGACCAAGAT	GCCCAACCAC	180
GGACAGGCGT	AGGGCGACAC	GTTTCAGGGA	GGGACCTTGT	ACTGGTTCTA	CGGGTTGGTG	
CTGCACCACA	GCACCCAGGC	TAACGCCATC	CTGGCCATGG	AACAGTTCGA	AGGGCTGCTG	240
GACGTGGTGT	CGTGGGTCCG	ATTGCGGTAG	GACCGGTACC	TTGTCAAGCT	TCCCGACGAC	
GGCACCCACT	GCAGCCCGGA	TCTTCTCTTC	TTCCTCTGTG	CAATGTACGC	ACCCATTTGC	300
CCGTGGGTGA	CGTCGGGCCCT	AGAAGAGAAG	AAGGAGACAC	GTTACATGCG	TGGGTAAACG	
ACCATCGACT	TCCAGCACGA	GCCCATCAAG	CCCTGCAAGT	CTGTGTGTGA	GCGCGCCCGA	360
TGGTAGCTGA	AGGTTCGTCT	CGGGTAGTTC	GGGACGTTCA	GACACACACT	CGCGCGGGCT	
CAGGGCTGCG	AGCCCATTTCT	CATCAAGTAC	CGCCACTCGT	GGCCGGAAAG	CTTGGCCTGC	420
GTCCCGACGC	TCGGGTAAAG	GTAGTTCATG	GCGGTGAGCA	CCGGCCTTTC	GAACCGGACG	
GACGAGCTGC	CGGTGTACGA	CCGCGGCGTG	TGCATCTCTC	CTGAGGCCAT	CGTCACCGCG	480
CTGCTCGACG	GCCACATGCT	GGCGCCGCAC	ACGTAGAGAG	GACTCCGGTA	GCAGTGGCGC	
GACGGAGCGG	ATTTTCCTAT	GGATTCAAGT	ACTGGACACT	GCAGAGGGGC	AAGCAGCGAA	540
CTGCCTCGCC	TAAAAGGATA	CCTAAGTTCA	TGACCTGTGA	CGTCTCCCCG	TTCGTCGCTT	
CGTTGCAAAAT	GTAAGCCTGT	CAGAGCTACA	CAGAAGACCT	ATTTCCGGAA	CAATTACAAC	600
GCAACGTTTA	CATTCGGACA	GTCTCGATGT	GTCTTCTGGA	TAAAGGCCTT	GTTAATGTTG	
TATGTCATCC	GGGCTAAAGT	TAAAGAGGTA	AAGATGAAAT	GTCATGATGT	GACCGCCGTT	660
ATACAGTAGG	CCCGATTTCA	ATTTCTCCAT	TTCTACTTTA	CAGTACTACA	CTGGCGGCAA	
GTGGAAGTGA	AGGAAATTCCT	AAAGGCATCA	CTGGTAAACA	TTCCAAGGGA	CACCGTCAAT	720
CACCTTCACT	TCCTTTAAGA	TTTCCGTAGT	GACCATTTGT	AAGGTTCCCT	GTGGCAGTTA	
CTTTATACCA	CCTCTGGCTG	CCTCTGTCCT	CCACTTACTG	TCAATGAGGA	ATATGTCATC	780
GAAATATGGT	GGAGACCGAC	GGAGACAGGA	GGTGAATGAC	AGTTACTCCT	TATACAGTAG	
ATGGGCTATG	AAGACGAGGA	ACGTTCCAGG	TTACTCTTGG	TAGAAGGCTC	TATAGCTGAG	840
TACCCGATAC	TTCTGCTCCT	TGCAAGGTCC	AATGAGAACC	ATCTTCCGAG	ATATCGACTC	
AAGTGGAAGG	ATCGGCTTGG	TAAGAAAGTC	AAGCGCTGGG	ATATGAAACT	CCGACACCTT	900
TTCACCTTCC	TAGCCGAACC	ATTCTTTCAG	TTCGCGACCC	TATACTTTGA	GGCTGTGGAA	
GGACTGGGTA	AAACTGATGC	TAGCGATTCC	ACTCAGAATC	AGAAGTCTGG	CAGGAACTCT	960
CCTGACCCAT	TTTGACTACG	ATCGCTAAGG	TGAGTCTTAG	TCTTCAGACC	GTCCTTGAGA	

AATCCCCGGC CAGCACGCAG CTAAATCCTG AAATGTAAAA GGCCACACCC ACGGACTCCC 1020
 TTAGGGGCGG GTCGTGCGTC GATTTAGGAC TTTACATTTT CCGGTGTGGG TGCCTGAGGG

 TTCTAAGACT GGCCTGGTG GACTAACAAA GGAAAACCGC ACAGTTGTGC TCGTGACCGA 1080
 AAGATTCTGA CCGCGACCAC CTGATTGTTT CTTTTTGGCG TGTCAACACG AGCACTGGCT

 TTGTTTACCG CAGACACCGC GTGGCTACCG AAGTTACTTC CGGTCCCCTT TCTCCTGCTT 1140
 AACAAATGGC GTCTGTGGCG CACCGATGGC TTCAATGAAG GCCAGGGGAA AGAGGACGAA

 CTTAATGGCG TGGGGTTAGA TCCTTTAATA TGTATATAT TCTGTTTCAT CAATCACGTG 1200
 GAATTACCGC ACCCCAATCT AGGAAATTAT ACAATATATA AGACAAAGTA GTTAGTGCAC

 GGGACTGTTT TTTTGCAACC AGAATAGTAA ATTAAATATG TTGATGCTAA GGTTCCTGTA 1260
 CCCTGACAAG AAAACGTTGG TCTTATCATT TAATTTATAC AACTACGATT CCAAAGACAT

 CTGGACTCCC TGGGTTTAAT TTGGTGTCT GTACCCTGAT TGAGAATGCA ATGTTTCATG 1320
 GACCTGAGGG ACCCAAATTA AACCACAAGA CATGGGACTA ACTCTTACGT TACAAAGTAC

 TAAAGAGAGA ATCCTGGTCA TATCTCAAGA ACTAGATATT GCTGTAAGAC AGCCTCTGCT 1380
 ATTTCTCTCT TAGGACCAGT ATAGAGTTCT TGATCTATAA CGACATTCTG TCGGAGACGA

 GCTGCGCTTA TAGTCTTG TGTTGTATGCC TTTGTCCATT TCCCTCATGC TGTGAAAGTT 1440
 CGACGCGAAT ATCAGAACAC AAACATACGG AAACAGGTAA AGGGAGTACG ACACTTTCAA

 ATACATGTTT ATAAAGGTAG AACGGCATT TGAAATCAGA CACTGCACAA GCAGAGTAGC 1500
 TATGTACAAA TATTTCCATC TTGCCGTAAA ACTTTAGTCT GTGACGTGTT CGTCTCATCG

 CCAACACCAG GAAGCATTTA TGAGGAAACG CCACACAGCA TGACTTATTT TCAAGATTGG 1560
 GGTGTGGTCT CTCGTAAAT ACTCCTTTGC GGTGTGTCGT ACTGAATAAA AGTTCTAACC

 CAGGCAGCAA AATAAATAGT GTTGGGAGCC AAGAAAAGAA TATTTTGCCT GGTAAAGGGG 1620
 GTCCGTCGTT TTATTTATCA CAACCCTCGG TTCTTTTCTT ATAAAACGGA CCAATTCCCC

 CACACTGGAA TCAGTAGCCC TTGAGCCATT AACAGCAGTG TTCTTCTGGC AAGTTTTTGA 1680
 GTGTGACCTT AGTCATCGGG AACTCGGTAA TTGTCGTCAC AAGAAGACCG TTCAAAAACCT

 TTTGTTTATA AATGTATTCA CGAGCATTAG AGATGAACTT ATAACTAGAC ATCTGTTGTT 1740
 AAACAAGTAT TTACATAAGT GCTCGTAATC TCTACTTGAA TATTGATCTG TAGACAACAA

 ATCTCTATAG CTCTGCTTCC TTCTAAATCA AACCCATTGT TGGATGCTCC CTCTCCATTC 1800
 TAGAGATATC GAGACGAAGG AAGATTTAGT TTGGGTAACA ACCTACGAGG GAGAGGTAAG

ATAAATAAAT	TTGGCTTGCT	GTATTGGCCA	GGAAAAGAAA	GTATTAAAGT	ATGCATGCAT	1860
TATTTATTTA	AACCGAACGA	CATAACCGGT	CCTTTTCTTT	CATAATTTCA	TACGTACGTA	
GTGCACCAGG	GTGTTATTTA	ACAGAGGTAT	GTAACCTCTAT	AAAAGACTAT	AATTTACAGG	1920
CACGTGGTCC	CACAATAAAT	TGTCTCCATA	CATTGAGATA	TTTTCTGATA	TTAAATGTCC	
ACACGGAAAT	GTGCACATTT	GTTTACTTTT	TTTCTTCCTT	TTGCTTTGGG	CTTGTGATTT	1980
TGTGCCTTTA	CACGTGTAAA	CAAATGAAAA	AAAGAAGGAA	AACGAAACCC	GAACACTAAA	
TGGTTTTTGG	TGTGTTTATG	TCTGTATTTT	GGGGGGTGGG	TAGGTTTAAG	CCATTGCACA	2040
ACAAAAAACC	ACACAAATAC	AGACATAAAA	CCCCCACC	ATCCAAATTC	GGTAACGTGT	
TTCAAGTTGA	ACTAGATTAG	AGTAGACTAG	GCTCATTGGC	CTAGACATTA	TGATTTGAAT	2100
AAGTTCAACT	TGATCTAATC	TCATCTGATC	CGAGTAACCG	GATCTGTAAT	ACTAACTTA	
TTGTGTTGTT	TAATGCTCCA	TCAAGATGTC	TAATAAAAGG	AATATGGTTG	TCAACAGAGA	2160
AACACAACAA	ATTACGAGGT	AGTTCTACAG	ATTATTTTCC	TTATACCAAC	AGTTGTCTCT	
CGACAACAAC	AACAAA					
GCTGTTGTTG	TTGTTT					

1860
 1920
 1980
 2040
 2100
 2160

MVCGSPGGML LLRAGLLALA ALCLLRVPGA RAAACEPVRI PLCKSLPWNM TKMPNHLHHS 60
TQANAILAIE QFEGLLGTHC SPDLLFFLCA MYAPICTIDF QHEPIKPCKS VCERARQGCE 120
PILIKYRHSW PENLACEELP VYDRGVCISP EAIVTADGAD FPMDSNGNC RGASSERCKC 180
KPIRATQKTY FRNNYNYVIR AKVKEIKTKC HDVTAVVEVK EILKSSLVNI PRDTVNLYTS 240
SGCLCPPLNV NEEYIIMGYE DEERSRLLLV EGSIAEKWKD RLGKKVKRWD MKLRHLGLSK 300
SDSSNSDSTQ SQKSGRNSNP RQARN.

Figure 9. Deduced amino acid sequence of human FRZB-1 protein. SEQ ID NO:9.

Figure 10. Nucleotide sequence of the full-length human FRZB-1 cDNA. SEQ ID NO:10.
This sequence was assembled from public ESTs from the Genbank database
(accession numbers: H18848, R63748, W38677, W44760, H38379 and N71244).

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GGCGGAGCGG GCCTTTTGGC GTCCACTGCG CGGCTGCACC CTGCCCCATC TGCCGGGATC      60
CCGCCTCGCC CGGAAAACCG CAGGTGACGC GCCGACGTGG GACGGGGTAG ACGGCCCTAG

ATGGTCTGCG GCAGCCCGGG AGGGATGCTG CTGCTGCGGG CCGGGCTGCT TGCCCTGGCT      120
TACCAGACGC CGTCGGGGCC TCCCTACGAC GACGACGCCC GGCCCGACGA ACGGGACCGA

GCTCTCTGCC TGCTCCGGGT GCCCGGGGCT CGGGCTGCAG CCTGTGAGCC CGTCCGCATC      180
CGAGAGACGG ACGAGGCCCA CGGGCCCCGA GCCCGACGTC GGACACTCGG GCAGGCGTAG

CCCCTGTGCA AGTCCCTGCC CTGGAACATG ACTAAGATGC CCAACCACCT GCACCACAGC      240
GGGGACACGT TCAGGGACGG GACCTTGTAC TGATTCTACG GGTGGGTGGA CGTGGTGTCTG

ACTCAGGCCA ACGCCATCCT GGCCATCGAG CAGTTCGAAG GTCTGCTGGG CACCCACTGC      300
TGAGTCCGGT TGCGGTAGGA CCGGTAGCTC GTCAAGCTTC CAGACGACCC GTGGGTGACG

AGCCCCGATC TGCTCTTCTT CCTCTGTGCC ATGTACGCGC CCATCTGCAC CATTGACTTC      360
TCGGGGCTAG ACGAGAAGAA GGAGACACGG TACATGCGCG GGTAGACGTG GTAAGTGAAG

CAGCACGAGC CCATCAAGCC CTGTAAGTCT GTGTGCGAGC GGGCCCCGCA GGGCTGTGAG      420
GTCGTGCTCG GGTAGTTCGG GACATTGAGA CACACGCTCG CCCGGGCCGT CCCGACACTC

CCCATACTCA TCAAGTACCG CCACTCGTGG CCGGAGAACC TGGCCTGCGA GGAGCTGCCA      480
GGGTATGAGT AGTTCATGGC GGTGAGCACC GGCCTCTTGG ACCGGACGCT CCTCGACGGT

GTGTACGACA GGGGCGTGTG CATCTCTCCC GAGGCCATCG TTAAGTGGGA CGGAGCTGAT      540
CACATGCTGT CCGCGCACAC GTAGAGAGGG CTCCGGTAGC AATGACGCCT GCCTCGACTA

TTTCCTATGG ATTCTAGTAA CGGAAACTGT AGAGGGGCAA GCAGTGAACG CTGTAAATGT      600
AAAGGATACC TAAGATCATT GCCTTTGACA TCTCCCCGTT CGTCACTTGC GACATTTACA

AAGCCTATTA GAGCTACACA GAAGACCTAT TTCCGGAACA ATTACAATA TGTCATTTCGG      660
TTCGGATAAT CTCGATGTGT CTTCTGGATA AAGGCCTTGT TAATGTTGAT ACAGTAAGCC

GCTAAAGTTA AAGAGATAAA GACTAAGTGC CATGATGTGA CTGCAGTAGT GGAGGTGAAG      720
CGATTTCAAT TTCTCTATTT CTGATTCACG GTACTACACT GACGTCATCA CCTCCACTTC

GAGATTCTAA AGTCCTCTCT GTTAAACATT CCACGGGACA CTGTCAACCT CTATACCAGC      780
CTCTAAGATT TCAGGAGAGA CCATTTGTAA GGTGCCCTGT GACAGTTGGA GATATGGTCG

TCTGGCTGCC TCTGCCCTCC ACTTAATGTT AATGAGGAAT ATATCATCAT GGGCTATGAA      840
AGACCGACGG AGACGGGAGG TGAATTACAA TTAAGCTTAA TATAGTAGTA CCCGATACTT

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